RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/579,693
Source: IFW
Date Processed by STIC: 5/31/06

ENTERED

IFWP

RAW SEQUENCE LISTING DATE: 05/31/2006
PATENT APPLICATION: **US/10/579,693** TIME: 13:29:04

Input Set : **E:\seqlist.txt**
Output Set: **N:\CRF4\05312006\J579693.raw**

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      4 <110> APPLICANT: BASF AKTIENGESELLSCHAFT et al.
      6 <120> TITLE OF INVENTION: METHODS FOR THE PREPARATION OF A FINE
      7       CHEMICAL BY FERMENTATION
     10 <130> FILE REFERENCE: BGI-159PC2
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/579,693
C--> 13 <141> CURRENT FILING DATE: 2006-05-18
     15 <150> PRIOR APPLICATION NUMBER: PCT/IB2003/006464
     16 <151> PRIOR FILING DATE: 2003-12-18
     18 <160> NUMBER OF SEQ ID NOS: 15
     20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     22 <210> SEQ ID NO: 1
     23 <211> LENGTH: 1650
     24 <212> TYPE: DNA
     25 <213> ORGANISM: Corynebacterium glutamicum
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     28 <221> NAME/KEY: CDS
     29 <222> LOCATION: (101)...(1627)
     31 <400> SEQUENCE: 1
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     33 aaacactgct tagtggccca atacgtgcaa aaataaggcc atg aga atc tca aag   115
     34                                              Met Arg Ile Ser Lys
     35                                               1               5
     37 gcc aat gcg tat gtt gca gcg att gac caa ggc acc act tcc act cgg   163
     38 Ala Asn Ala Tyr Val Ala Ala Ile Asp Gln Gly Thr Thr Ser Thr Arg
     39                  10                  15                  20
     41 tgc atc ttc att gat gcc caa gga aaa gtg gtg tct tct gct tcc aag   211
     42 Cys Ile Phe Ile Asp Ala Gln Gly Lys Val Val Ser Ser Ala Ser Lys
     43              25                  30                  35
     45 gag cac cgc caa atc ttc cca caa cag ggc tgg gta gag cac gat cct   259
     46 Glu His Arg Gln Ile Phe Pro Gln Gln Gly Trp Val Glu His Asp Pro
     47          40                  45                  50
     49 gaa gaa att tgg gac aac att cga tct gtc gtc agc cag gcg atg gtc   307
     50 Glu Glu Ile Trp Asp Asn Ile Arg Ser Val Val Ser Gln Ala Met Val
     51      55                  60                  65
     53 tcc att gac atc acc cca cac gag gtt gca tcg ctg gga gtc acc aac   355
     54 Ser Ile Asp Ile Thr Pro His Glu Val Ala Ser Leu Gly Val Thr Asn
     55  70                  75                  80                  85
     57 cag cgc gaa acc acc gtg gtg tgg gac aag cac acc ggc gaa cct gtc   403
     58 Gln Arg Glu Thr Thr Val Val Trp Asp Lys His Thr Gly Glu Pro Val
     59                  90                  95                 100
     61 tac aac gca atc gtg tgg caa gac acc cgc acc tct gac att tgc cta   451
     62 Tyr Asn Ala Ile Val Trp Gln Asp Thr Arg Thr Ser Asp Ile Cys Leu
     63             105                 110                 115
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65 gag atc gcg ggc gaa gaa ggc cag gaa aag tgg ctt gac cgc acc ggc   499
66 Glu Ile Ala Gly Glu Glu Gly Gln Glu Lys Trp Leu Asp Arg Thr Gly
67         120                 125                 130
69 ctg ctg atc aac tcc tac cca tcg ggg ccc aaa atc aag tgg att ctc   547
70 Leu Leu Ile Asn Ser Tyr Pro Ser Gly Pro Lys Ile Lys Trp Ile Leu
71     135                 140                 145
73 gac aac gtt gag gga gct cgc gaa cgc gcc gaa aag ggc gac ctt ttg   595
74 Asp Asn Val Glu Gly Ala Arg Glu Arg Ala Glu Lys Gly Asp Leu Leu
75 150                 155                 160                 165
77 ttt ggc acc atg gat acc tgg gtg ctg tgg aac ctg acc ggc ggt gtc   643
78 Phe Gly Thr Met Asp Thr Trp Val Leu Trp Asn Leu Thr Gly Gly Val
79                 170                 175                 180
81 cgc ggc gac gac ggt gat gat gcc atc cac gtc acc gat gtc acc aac   691
82 Arg Gly Asp Asp Gly Asp Asp Ala Ile His Val Thr Asp Val Thr Asn
83             185                 190                 195
85 gca tcc cgc aca cta ttg atg gat ctc cgc acg caa cag tgg gat cca   739
86 Ala Ser Arg Thr Leu Leu Met Asp Leu Arg Thr Gln Gln Trp Asp Pro
87         200                 205                 210
89 gaa cta tgc gaa gcc cta gac att ccg atg tcc atg ctc cct gag atc   787
90 Glu Leu Cys Glu Ala Leu Asp Ile Pro Met Ser Met Leu Pro Glu Ile
91     215                 220                 225
93 cgt ccc tcc gtc gga gaa ttc cgc tcc gtg cgc cac cgc gga acc cta   835
94 Arg Pro Ser Val Gly Glu Phe Arg Ser Val Arg His Arg Gly Thr Leu
95 230                 235                 240                 245
97 gcc gac gtc ccg att act ggc gtg ctc ggc gac cag caa gcg gcc ctt   883
98 Ala Asp Val Pro Ile Thr Gly Val Leu Gly Asp Gln Gln Ala Ala Leu
99                 250                 255                 260
101 ttt ggt cag ggc gga ttc cac gaa ggt gct gct aaa aat acc tac ggc   931
102 Phe Gly Gln Gly Gly Phe His Glu Gly Ala Ala Lys Asn Thr Tyr Gly
103             265                 270                 275
105 acc ggc ctc ttc ctg ctg atg aac acc ggc acc tcg ttg aag att tcc   979
106 Thr Gly Leu Phe Leu Leu Met Asn Thr Gly Thr Ser Leu Lys Ile Ser
107         280                 285                 290
109 gag cac ggc ctg ctg tcc acc atc gcc tat caa cgg gaa gga tcc gct   1027
110 Glu His Gly Leu Leu Ser Thr Ile Ala Tyr Gln Arg Glu Gly Ser Ala
111     295                 300                 305
113 ccg gtc tac gcg ctg gaa ggt tcc gta tcc atg ggc ggt tcc ttg gtg   1075
114 Pro Val Tyr Ala Leu Glu Gly Ser Val Ser Met Gly Gly Ser Leu Val
115 310                 315                 320                 325
117 cag tgg ctg cgc gac aac cta cag cta atc ccc aac gca cca gcg att   1123
118 Gln Trp Leu Arg Asp Asn Leu Gln Leu Ile Pro Asn Ala Pro Ala Ile
119                 330                 335                 340
121 gaa aac ctc gcc cga gaa gtc gaa gac aac ggt ggc gtt cat gtt gtc   1171
122 Glu Asn Leu Ala Arg Glu Val Glu Asp Asn Gly Gly Val His Val Val
123             345                 350                 355
125 cca gca ttc acc gga ctg ttc gca cca cgt tgg cgc ccc gat gct cgt   1219
126 Pro Ala Phe Thr Gly Leu Phe Ala Pro Arg Trp Arg Pro Asp Ala Arg
127         360                 365                 370
129 ggc gtc att aca ggc ctc acc cgt ttt gcc aac cgc aaa cac atc gcc   1267
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130 Gly Val Ile Thr Gly Leu Thr Arg Phe Ala Asn Arg Lys His Ile Ala
131     375                 380                 385
133 cgc gca gtc ctt gaa gcc aac gcc ttc caa acc cgc gaa gtt gtg gac   1315
134 Arg Ala Val Leu Glu Ala Asn Ala Phe Gln Thr Arg Glu Val Val Asp
135 390                 395                 400                 405
137 gcc atg gcc aaa gac gca ggc aaa gcc ctc gaa tcc ctc cgc gtc gac   1363
138 Ala Met Ala Lys Asp Ala Gly Lys Ala Leu Glu Ser Leu Arg Val Asp
139                 410                 415                 420
141 ggt gcg atg gtg gaa aat gac ctc ctc atg caa atg caa gcc gac ttc   1411
142 Gly Ala Met Val Glu Asn Asp Leu Leu Met Gln Met Gln Ala Asp Phe
143             425                 430                 435
145 ctc ggc atc gac gtc caa cgt ctc gag gac gta gaa acc acc gcc gtc   1459
146 Leu Gly Ile Asp Val Gln Arg Leu Glu Asp Val Glu Thr Thr Ala Val
147         440                 445                 450
149 ggc gtc gca ttc gct gca ggt ctc ggc tct gga ttc ttc aaa aca act   1507
150 Gly Val Ala Phe Ala Ala Gly Leu Gly Ser Gly Phe Phe Lys Thr Thr
151     455                 460                 465
153 gac gag atc gaa aaa ctt att gca gtg aag aaa gtc tgg aac cct gac   1555
154 Asp Glu Ile Glu Lys Leu Ile Ala Val Lys Lys Val Trp Asn Pro Asp
155 470                 475                 480                 485
157 atg agc gaa gaa gag cgc gaa cgt cgc tat gcc gaa tgg aat agg gca   1603
158 Met Ser Glu Glu Glu Arg Glu Arg Arg Tyr Ala Glu Trp Asn Arg Ala
159                 490                 495                 500
161 gtg gag cat tct tat gac cag gcc tagctgattt gggtcggcct tta         1650
162 Val Glu His Ser Tyr Asp Gln Ala
163             505
166 <210> SEQ ID NO: 2
167 <211> LENGTH: 509
168 <212> TYPE: PRT
169 <213> ORGANISM: Corynebacterium glutamicum
171 <400> SEQUENCE: 2
172 Met Arg Ile Ser Lys Ala Asn Ala Tyr Val Ala Ala Ile Asp Gln Gly
173  1               5                  10                  15
174 Thr Thr Ser Thr Arg Cys Ile Phe Ile Asp Ala Gln Gly Lys Val Val
175             20                  25                  30
176 Ser Ser Ala Ser Lys Glu His Arg Gln Ile Phe Pro Gln Gln Gly Trp
177         35                  40                  45
178 Val Glu His Asp Pro Glu Glu Ile Trp Asp Asn Ile Arg Ser Val Val
179     50                  55                  60
180 Ser Gln Ala Met Val Ser Ile Asp Ile Thr Pro His Glu Val Ala Ser
181 65                  70                  75                  80
182 Leu Gly Val Thr Asn Gln Arg Glu Thr Thr Val Val Trp Asp Lys His
183                 85                  90                  95
184 Thr Gly Glu Pro Val Tyr Asn Ala Ile Val Trp Gln Asp Thr Arg Thr
185             100                 105                 110
186 Ser Asp Ile Cys Leu Glu Ile Ala Gly Glu Glu Gly Gln Glu Lys Trp
187         115                 120                 125
188 Leu Asp Arg Thr Gly Leu Leu Ile Asn Ser Tyr Pro Ser Gly Pro Lys
189     130                 135                 140
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PATENT APPLICATION: **US/10/579,693** TIME: 13:29:04

Input Set : **E:\seqlist.txt**
Output Set: **N:\CRF4\05312006\J579693.raw**

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190 Ile Lys Trp Ile Leu Asp Asn Val Glu Gly Ala Arg Glu Arg Ala Glu
191 145                 150                 155                 160
192 Lys Gly Asp Leu Leu Phe Gly Thr Met Asp Thr Trp Val Leu Trp Asn
193                 165                 170                 175
194 Leu Thr Gly Gly Val Arg Gly Asp Asp Gly Asp Asp Ala Ile His Val
195             180                 185                 190
196 Thr Asp Val Thr Asn Ala Ser Arg Thr Leu Leu Met Asp Leu Arg Thr
197         195                 200                 205
198 Gln Gln Trp Asp Pro Glu Leu Cys Glu Ala Leu Asp Ile Pro Met Ser
199     210                 215                 220
200 Met Leu Pro Glu Ile Arg Pro Ser Val Gly Glu Phe Arg Ser Val Arg
201 225                 230                 235                 240
202 His Arg Gly Thr Leu Ala Asp Val Pro Ile Thr Gly Val Leu Gly Asp
203                 245                 250                 255
204 Gln Gln Ala Ala Leu Phe Gly Gln Gly Gly Phe His Glu Gly Ala Ala
205             260                 265                 270
206 Lys Asn Thr Tyr Gly Thr Gly Leu Phe Leu Leu Met Asn Thr Gly Thr
207         275                 280                 285
208 Ser Leu Lys Ile Ser Glu His Gly Leu Leu Ser Thr Ile Ala Tyr Gln
209     290                 295                 300
210 Arg Glu Gly Ser Ala Pro Val Tyr Ala Leu Glu Gly Ser Val Ser Met
211 305                 310                 315                 320
212 Gly Gly Ser Leu Val Gln Trp Leu Arg Asp Asn Leu Gln Leu Ile Pro
213                 325                 330                 335
214 Asn Ala Pro Ala Ile Glu Asn Leu Ala Arg Glu Val Glu Asp Asn Gly
215             340                 345                 350
216 Gly Val His Val Val Pro Ala Phe Thr Gly Leu Phe Ala Pro Arg Trp
217         355                 360                 365
218 Arg Pro Asp Ala Arg Gly Val Ile Thr Gly Leu Thr Arg Phe Ala Asn
219     370                 375                 380
220 Arg Lys His Ile Ala Arg Ala Val Leu Glu Ala Asn Ala Phe Gln Thr
221 385                 390                 395                 400
222 Arg Glu Val Val Asp Ala Met Ala Lys Asp Ala Gly Lys Ala Leu Glu
223                 405                 410                 415
224 Ser Leu Arg Val Asp Gly Ala Met Val Glu Asn Asp Leu Leu Met Gln
225             420                 425                 430
226 Met Gln Ala Asp Phe Leu Gly Ile Asp Val Gln Arg Leu Glu Asp Val
227         435                 440                 445
228 Glu Thr Thr Ala Val Gly Val Ala Phe Ala Ala Gly Leu Gly Ser Gly
229     450                 455                 460
230 Phe Phe Lys Thr Thr Asp Glu Ile Glu Lys Leu Ile Ala Val Lys Lys
231 465                 470                 475                 480
232 Val Trp Asn Pro Asp Met Ser Glu Glu Glu Arg Glu Arg Arg Tyr Ala
233                 485                 490                 495
234 Glu Trp Asn Arg Ala Val Glu His Ser Tyr Asp Gln Ala
235             500                 505
238 <210> SEQ ID NO: 3
239 <211> LENGTH: 35
240 <212> TYPE: DNA
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PATENT APPLICATION: **US/10/579,693** TIME: 13:29:04

Input Set : **E:\seqlist.txt**
Output Set: **N:\CRF4\05312006\J579693.raw**

<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Oligonucleotide
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<211> LENGTH: 34
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Oligonucleotide
<400> SEQUENCE: 4
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<210> SEQ ID NO: 5
<211> LENGTH: 4323
<212> TYPE: DNA
<213> ORGANISM: Corynebacterium glutamicum
<400> SEQUENCE: 5
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VERIFICATION SUMMARY DATE: 05/31/2006
PATENT APPLICATION: **US/10/579,693** TIME: 13:29:05

Input Set : **E:\seqlist.txt**
Output Set: **N:\CRF4\05312006\J579693.raw**

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date